4. (Original) A constructed DNA sequence including a protein coding region encoding a modified barley α-glucosidase enzyme, the modified barley α-glucosidase differing from the wild-type barley α-glucosidase by the presence of a proline residue at residue 340.

5. (Original) A transgenic host which expresses the constructed DNA sequence of claim 4.

6. (Original) A modified α-glucosidase enzyme, the modified form differing from the wild-type barley α-glucosidase by at least one amino acid substitution from the native barley α-glucosidase enzyme sequence, the modified enzyme retaining enzymatic activity at a higher temperature than the wild-type enzyme.

7. (Amended) A modified α-glucosidase enzyme as claimed in claim 6 wherein the modification is selected from the group consisting of ~~removing an aspartate at residue 83, removing an aspartate from residue 92, adding a proline to residue 100,~~ adding a proline and removing an aspartate at residue 101, removing an aspartate from residue 105, ~~adding a proline to residue 122, adding a proline to residue 184, removing N-glycosylation site from residue 298, adding a proline to residue 336,~~ removing an aspartate from residue 369, adding N-glycosylation site and removing an aspartate from residue 372, ~~removing N-glycosylation site from residue 391, adding a proline to residue 394, adding a proline and removing an aspartate at residue 403,~~ adding N-glycosylation site to residue 463, removing an aspartate from residue 508, ~~removing a deamidation site from residue 568,~~ adding N-glycosylation site and removing an aspartate from residue 694, ~~adding a proline at residue 713, adding a proline at residue 742,~~ and removing an aspartate from residue 764.

8. (Original) A DNA sequence which encodes the modified α-glucosidase enzyme as claimed in claim 7.

9. (Amended) A method of making a mutant form of the enzyme barley α-glucosidase comprising the steps of:

(a) constructing a mutant gene sequence encoding a mutant form of the α-glucosidase enzyme;

(b) cloning the mutant gene sequence into an expression vector;

(c) expressing the protein encoded by the expression vector to produce the protein encoded by the mutant gene sequence;

(d) recovering the protein produced; and

(e) testing the protein for both α-glucosidase activity and for thermostability; wherein the mutant gene sequence encoding a mutant protein has at least one mutation selected from the group consisting of ~~removing an aspartate at residue 83, removing an aspartate from residue 92, adding a proline to residue 100,~~ adding a proline and removing an aspartate at residue 101, removing an aspartate from residue 105, ~~adding a proline to residue 122, adding a proline to residue 184, removing N-glycosylation site from residue 298, adding a proline to residue 336,~~ removing an aspartate from residue 369, adding N-glycosylation site and removing an aspartate from residue 372, ~~removing N-glycosylation site from residue 391, adding a proline to residue 394, adding a proline and removing an aspartate at residue 403,~~ adding N-glycosylation site to residue 463, removing an aspartate from residue 508, ~~removing a deamidation site from residue 568,~~ adding N-glycosylation site and removing an aspartate from residue 694, ~~adding a proline at residue 713, adding a proline at residue 742,~~ and removing an aspartate from residue 764.